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Amendments to the Claims

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- 1. (Currently Amended) A scalable process for the highly selective, high yield separation of nucleic acids, comprising in combination: exposure introduction, enhancement, or stabilization of structural "affinity handles" from shielded purine base sites previously present, by a process selected from the group consisting of: selective thermal denaturation and renaturation, alkaline denaturation or the use of restriction enzymes yielding single-stranded overhangs, selectively to either the desired or the undesired moieties or nucleic acid; followed by capture of the undesired (or desired) nucleic acids by techniques which are selective for the characteristics of the affinity handle wherein the product is either a nucleic acid to be purified, or a nucleic acid or a non-nucleic acid product, from which undesired nucleic acid is to be separated.
- 2. (Previously Presented) A process according to Claim 1 wherein the handle comprises a structural form selected from the group consisting of single stranded region of nucleic acid, Triplexes, Hairpins, Stems, Loops, Cruciforms, G quartets, and modifications to the phosphate backbone.
- 3. (Currently Amended) A process according to Claim 1 wherein the captured nucleic acid product comprises a moiety that is sensitive to host genomic DNA contamination during selective separation such as single-strandedness in the undesired (or desired) nucleic acids as compared to the usual structure, such as double-strandedness, of the desired (or undesired) nucleic acids.
- 4. (Currently Amended) A process according to Claim 1 wherein the captured nucleic acid product comprises single-strandedness in the desired product or a moeity that is

sensitive to host genomic DNA contamination selective as compared to the structure of the undesired nucleic acids.

- 5. (Currently Amended) A process according to Claim 1 [[4]] comprising manufacture of recombinant *Taq* polymerase.
- 6. (Currently Amended) A process according to Claim 4 wherein the exposed bases of single-stranded undesired (or desired) nucleic acids facilitate a separation step selected from the group comprising: immobilized metal affinity chromatography (IMAC), immobilized single-stranded DNA binding (SSB) protein, the use of immobilized nucleic acids (poly-dT, poly-dU, or specific sequences), and the use of peptide nucleic acids (PNAs).
- 7. (Previously Presented) A process according to Claim 1 comprising introducing single strandedness as a handle.
- 8. (Currently Amended) A process according to Claim 1 occurring after another a thermally based process (such as heat based microbial lysis), in which a nucleic acid contaminant (such as genomic DNA) is rapidly cooled to below 65°C and is captured by an affinity method.
- 9. [Currently Amended) A process according to Claim 1 performed after another <u>an</u> alkali based process (such as alkaline lysis), in which genomic DNA or other nucleic acid contaminant [[)]] is rapidly neutralized and is captured by an affinity method.

- 10. (Currently Amended) A process according to Claim 1 comprising a step for introducing handles selected from the group comprising: selective thermal denaturation and renaturation, alkaline denaturation, the use of chaotropic agents, the use of restriction enzymes yielding single-stranded overhangs, the use of oligonucleotide dTs, single-stranded DNA binding proteins, minerals, and the use of oligonucleotide dTs, single-stranded DNA binding proteins, minerals, primers or other nucleic acid fragments such as complementary DNA nucleic acids to facilitate capture and separation of the undesired (or desired) nucleic acid from the desired (or undesired) nucleic acids.
- 11. (Previously Presented) A process according to Claim 1 wherein other plasmid isoforms selected from the group consisting of open circular ("nicked") and linear are selectively removed from the desired supercoiled plasmid DNA.
- 12. (Previously Presented) A process according to Claim 9 wherein other plasmid isoforms selected from the group consisting of open circular and linear are selectively removed from supercoiled plasmid DNA.
- 14. (Previously Presented) A process according to Claim 1 in which the separation is achieved by adsorption on chelated metal.
- 15. (Previously Presented) A process according to Claim 1 in which the separation is achieved using multi-channel plates.
- 16. (Previously Presented) A process according to Claim 1 in which the separation is achieved using IMAC spin columns.
- 17. (Previously Presented) A process according to Claim 1 in which the separation is achieved using magnetic particles.

- 18. (Previously Presented) A process according to Claim 1 in which the separation of multiple samples is achieved in parallel fashion:
- 19. (Currently Amended) A process according to Claim 1 in which the captured nucleic acid product comprises a moiety selected from BACs, PACs and YACs protein.

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- 20. (Currently Amended) A process according to Claim 1 in which the desired captured nucleic acid product is a plasmid.
- 21. (Currently Amended) A process according to Claim 1 in which the desired captured nucleic acid product is genomic DNA.
- 22. (Currently Amended) A process according to Claim 1 in which the desired captured nucleic acid product is RNA.
- 23. (Currently Amended) A process according to Claim 1 in which the capture method comprises is HIC.
- 24. (Currently Amended) A process according to Claim 1 in which the capture method comprises is RPC.
- 25. Cancelled